FIGURE 1

A1:			
hβ1A		MGRLLALVVGAALVSSACGGCVEVDSETEAVYGMTFKILCISCKRRSETN	
rβ1A		MGTLLALVVGAVLVSSAWGGCVEVDSETEAVYGMTFKILCISCKRRSETT	
hβ1A		ABTFTEWTFRQKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKD	
rβ1A		ABTFTEWTFRQKGTEEFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKD	
•		LQDLSIFITNVTYNHSGDYECHVYRLLFFENYEHNTSVVKKIHIEVVDKG	
		LQDLSIFITNVTYNHSGDYECHVYRLLFFDNYEHNTSVVKKIHLEVVDKG	
•		ESGAACPFTVTHRRARWRDRWQAVDRTGWLCAWPANRPQQRAEGEGSSPS . .	
		$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	
•		CPLQLWPLFLSSPRRGQ.SMPVPHRRSGYRTQLCHLCCMTSGRCL.LSLS : : :	
		LVLETSALQHTGGQIRTPTPPPTNGMCIGL.HSCCVTSDGCIPISEP	237
hβ1A	249	QRVVLGLPGIIRCVSRGVV	
rβ1A B:	238	QACPQGPERIFCMACCVSQAGPHWRDVGTYLRPQWE 273	
β1Α	1	MGRLLALVVGAALVSSACGGCVEVDSETEAVYGMTFKILCISCKRRSETN	50
hβ1	1	MGRLLALVVGAALVSSACGGCVEVDSETEAVYGMTFKILCISCKRRSETN	50
hβ1A	51	AETFTEWTFRQKGTEBFVKILRYENEVLQLEEDERFEGRVVWNGSRGTKD	100
hß1	51	AETFTEWTFRQKGTEEFVKILRYENEVLQLBEDERFEGRVVWNGSRGTKD	100
hβ1A	101	LQDLSIFITNVTYNHSGDYECHVYRLLFFENYEHNTSVVKKIHIEVVDKG	150
hβ1	101	LQDLSIFITNVTYNHSGDYECHVYRLLFFENYEHNTSVVKKIHIEVVDKA	150
hβ1A	151	ESGAACPFTVTHRRARWRDRWQAVDRTGWLCAWPANRPQQ.RAEGEGSSP	199
hβ1	151	NRDMASIVSEIMMYVLIVVLTIWLVAEMIYCYKKIAAATETAAQ	194
hβ1A	200	SCPLQLWPLFLSSPRRGQSMPVPHRRSGYRTQLCHLCCMTSGRCLLSLSQ	249
hβ1	195	ENASEYLAITSESKENCTGVQVAE	218

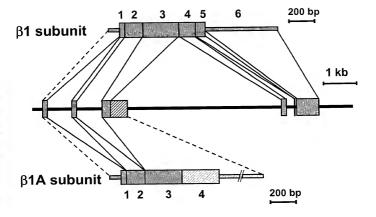


Figure 2

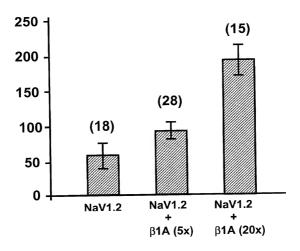
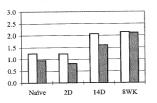
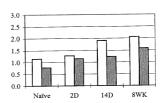


Figure 3

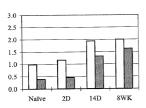
A. L5, ipsilateral DRGs



B. L5, contralateral DRGs



C. L4, ipsilateral DRGs



D. L4, contralateral DRGs

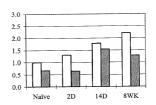
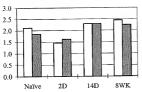
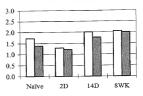


Figure 4

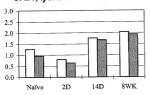
A. L5, ipsilateral DRGs



B. L5, contralateral DRGs



C. L4, ipsilateral DRGs



D. L4, contralateral DRGs

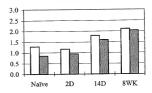


Figure 5